RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 09/857,581C
Source: IFW16
Date Processed by STIC: 6/27/06

ENTERED

CRF Errors Edited by the STIC Systems Branch

Serial Number: 09/857,581C CRF Edit Date: 6/27/06
Edited by: RV

___ Realigned nucleic acid/amino acid numbers/text in cases where the sequence text "wrapped" to the next line

___ Corrected the SEQ ID NO. Sequence numbers edited were:

_____

___ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

_____

___ Deleted: ___ invalid beginning/end-of-file text ; ___ page numbers

___ Inserted mandatory headings/numeric identifiers, specifically:

_____

___ Moved responses to same line as heading/numeric identifier, specifically:

_____

___ Other:

_____
_____
_____

Revised 09/09/2003

IFW16

RAW SEQUENCE LISTING DATE: 06/27/2006
PATENT APPLICATION: **US/09/857,581C** TIME: 16:52:46

Input Set : **A:\PTO.AMC.txt**
Output Set: **N:\CRF4\06272006\I857581C.raw**

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      3 <110> APPLICANT: Fader, Gary M.
      4       Jung, Woosuk
      5       Brian, McGonigle
      6       Odell, Joan T.
      7       Yu, Xiaodan
      9 <120> TITLE OF INVENTION: Nucleic Acid Fragments Encoding Isoflavone Synthase
     11 <130> FILE REFERENCE: BB1339RCE
     13 <140> CURRENT APPLICATION NUMBER: 09/857,581C
C--> 14 <141> CURRENT FILING DATE: 2001-06-05
     16 <150> PRIOR APPLICATION NUMBER: PCT/US00/01772
     17 <151> PRIOR FILING DATE: 2000-01-26
     19 <150> PRIOR APPLICATION NUMBER: 60/117,769
     20 <151> PRIOR FILING DATE: 1999-01-27
     22 <150> PRIOR APPLICATION NUMBER: 60/144,783
     23 <151> PRIOR FILING DATE: 1990-07-20
     25 <150> PRIOR APPLICATION NUMBER: 60/156,094
     26 <151> PRIOR FILING DATE: 1999-09-24
     28 <160> NUMBER OF SEQ ID NOS: 66
     30 <170> SOFTWARE: PatentIn version 3.3
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     34 <212> TYPE: DNA
     35 <213> ORGANISM: Glycine max
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     40 cgtcccacac caagtgcaaa atcaaaagca cttcgccacc tcccaaaccc tccaagccca    180
     41 aagcctcgtc ttcccttcat tggccacctt cacctcttaa aagataaact tctccactat    240
     42 gcactcatcg atctctccaa aaagcatggc cccttattct ctctctcctt cggctccatg    300
     43 ccaaccgtcg ttgcctccac ccctgagttg ttcaagctct tcctccaaac ccacgaggca    360
     44 acttccttca acacaaggtt ccaaacctct gccataagac gcctcactta cgacaactct    420
     45 gtggccatgg ttccattcgg accttactgg aagttcgtga ggaagctcat catgaacgac    480
     46 cttctcaacg ccaccaccgt caacaagctc aggcctttga ggacccaaca gatccgcaag    540
     47 ttccttaggg ttatggccca aagcgcagag gcccagaagc cccttgacgt caccgaggag    600
     48 cttctcaaat ggaccaacag caccatctcc atgatgatgc tcggcgaggc tgaggagatc    660
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     55 gagctcatca acaatcccag ggtgttgcaa aaggctcgtg aggaggtcta cagtgttgtg   1080
     56 ggcaaagata gactcgttga cgaagttgac actcaaaacc ttccttacat tagggccatt   1140
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RAW SEQUENCE LISTING DATE: 06/27/2006
PATENT APPLICATION: **US/09/857,581C** TIME: 16:52:46

Input Set : **A:\PTO.AMC.txt**
Output Set: **N:\CRF4\06272006\I857581C.raw**

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59 caagtaggaa gggaccccaa atactgggac agaccatcag aattccgtcc cgagaggttc 1320
60 ttagaaactg gtgctgaagg ggaagcaggg cctcttgatc ttaggggcca gcatttccaa 1380
61 ctcctcccat ttgggtctgg gaggagaatg tgccctggtg tcaatttggc tacttcagga 1440
62 atggcaacac ttcttgcatc tcttatccaa tgctttgacc tgcaagtgct gggccctcaa 1500
63 ggacaaatat tgaaaggtga tgatgccaaa gttagcatgg aagagagagc tggcctcaca 1560
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65 ctccttttctt aattaagata atcatcatat acaatagtag tgtcttgcca tcgcagttgc 1680
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70 <211> LENGTH: 521
71 <212> TYPE: PRT
72 <213> ORGANISM: Glycine max
74 <400> SEQUENCE: 2
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79 20 25 30
81 Pro Asn Pro Pro Ser Pro Lys Pro Arg Leu Pro Phe Ile Gly His Leu
82 35 40 45
84 His Leu Leu Lys Asp Lys Leu Leu His Tyr Ala Leu Ile Asp Leu Ser
85 50 55 60
87 Lys Lys His Gly Pro Leu Phe Ser Leu Ser Phe Gly Ser Met Pro Thr
88 65 70 75 80
90 Val Val Ala Ser Thr Pro Glu Leu Phe Lys Leu Phe Leu Gln Thr His
91 85 90 95
93 Glu Ala Thr Ser Phe Asn Thr Arg Phe Gln Thr Ser Ala Ile Arg Arg
94 100 105 110
96 Leu Thr Tyr Asp Asn Ser Val Ala Met Val Pro Phe Gly Pro Tyr Trp
97 115 120 125
99 Lys Phe Val Arg Lys Leu Ile Met Asn Asp Leu Leu Asn Ala Thr Thr
100 130 135 140
102 Val Asn Lys Leu Arg Pro Leu Arg Thr Gln Gln Ile Arg Lys Phe Leu
103 145 150 155 160
105 Arg Val Met Ala Gln Ser Ala Glu Ala Gln Lys Pro Leu Asp Val Thr
106 165 170 175
108 Glu Glu Leu Leu Lys Trp Thr Asn Ser Thr Ile Ser Met Met Met Leu
109 180 185 190
111 Gly Glu Ala Glu Glu Ile Arg Asp Ile Ala Arg Glu Val Leu Lys Ile
112 195 200 205
114 Phe Gly Glu Tyr Ser Leu Thr Asp Phe Ile Trp Pro Leu Lys Tyr Leu
115 210 215 220
117 Lys Val Gly Lys Tyr Glu Lys Arg Ile Asp Asp Ile Leu Asn Lys Phe
118 225 230 235 240
120 Asp Pro Val Val Glu Arg Val Ile Lys Lys Arg Arg Glu Ile Val Arg
121 245 250 255
123 Arg Arg Lys Asn Gly Glu Val Val Glu Gly Glu Ala Ser Gly Val Phe

RAW SEQUENCE LISTING DATE: 06/27/2006
PATENT APPLICATION: US/09/857,581C TIME: 16:52:46

Input Set : A:\PTO.AMC.txt
Output Set: N:\CRF4\06272006\I857581C.raw

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126 Leu Asp Thr Leu Leu Glu Phe Ala Glu Asp Glu Thr Met Glu Ile Lys
127          275                 280                 285
129 Ile Thr Lys Glu Gln Ile Lys Gly Leu Val Val Asp Phe Phe Ser Ala
130      290                 295                 300
132 Gly Thr Asp Ser Thr Ala Val Ala Thr Glu Trp Ala Leu Ala Glu Leu
133 305                 310                 315                 320
135 Ile Asn Asn Pro Arg Val Leu Gln Lys Ala Arg Glu Glu Val Tyr Ser
136                 325                 330                 335
138 Val Val Gly Lys Asp Arg Leu Val Asp Glu Val Asp Thr Gln Asn Leu
139              340                 345                 350
141 Pro Tyr Ile Arg Ala Ile Val Lys Glu Thr Phe Arg Met His Pro Pro
142          355                 360                 365
144 Leu Pro Val Val Lys Arg Lys Cys Thr Glu Glu Cys Glu Ile Asn Gly
145      370                 375                 380
147 Tyr Val Ile Pro Glu Gly Ala Leu Val Leu Phe Asn Val Trp Gln Val
148 385                 390                 395                 400
150 Gly Arg Asp Pro Lys Tyr Trp Asp Arg Pro Ser Glu Phe Arg Pro Glu
151                 405                 410                 415
153 Arg Phe Leu Glu Thr Gly Ala Glu Gly Glu Ala Gly Pro Leu Asp Leu
154              420                 425                 430
156 Arg Gly Gln His Phe Gln Leu Leu Pro Phe Gly Ser Gly Arg Arg Met
157          435                 440                 445
159 Cys Pro Gly Val Asn Leu Ala Thr Ser Gly Met Ala Thr Leu Leu Ala
160      450                 455                 460
162 Ser Leu Ile Gln Cys Phe Asp Leu Gln Val Leu Gly Pro Gln Gly Gln
163 465                 470                 475                 480
165 Ile Leu Lys Gly Asp Asp Ala Lys Val Ser Met Glu Glu Arg Ala Gly
166                 485                 490                 495
168 Leu Thr Val Pro Arg Ala His Ser Leu Val Cys Val Pro Leu Ala Arg
169              500                 505                 510
171 Ile Gly Val Ala Ser Lys Leu Leu Ser
172          515                 520
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176 <212> TYPE: DNA
177 <213> ORGANISM: Artificial Sequence
179 <220> FEATURE:
180 <223> OTHER INFORMATION: Oligonucleotide primer used in construction of WHT1
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188 <213> ORGANISM: Artificial Sequence
190 <220> FEATURE:
191 <223> OTHER INFORMATION: Oligonucleotide primer used in construction of yeast strain
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193 <400> SEQUENCE: 4
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RAW SEQUENCE LISTING DATE: 06/27/2006
PATENT APPLICATION: **US/09/857,581C** TIME: 16:52:46

Input Set : **A:\PTO.AMC.txt**
Output Set: **N:\CRF4\06272006\I857581C.raw**

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<220> FEATURE:
<223> OTHER INFORMATION: Oligonucleotide primer
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<210> SEQ ID NO: 8
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<213> ORGANISM: Artificial Sequence
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ctgcaaaatc aaaagcactt cgccatctcc caaacccacc aagcccaaag cctcgtcttc 180
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cctccacacc agaattgttc aagctcttcc tccaaacgca cgaggcaact tccttcaaca 360
caaggttcca aacctcagcc ataagacgcc tcacctatga tagctcagtg gccatggttc 420
ccttcggacc ttactggaag ttcgtgagga agctcatcat gaacgacctt cccaacgcca 480
ccactgtaaa caagttgagg cctttgagga cccaacagac ccgcaagttc cttagggtta 540
tggcccaagg cgcagaggca cagaagcccc ttgacttgac cgaggagctt ctgaaatgga 600
ccaacagcac catctccatg atgatgctcg gcgaggctga ggagatcaga gacatcgctc 660
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RAW SEQUENCE LISTING DATE: 06/27/2006
PATENT APPLICATION: **US/09/857,581C** TIME: 16:52:46

Input Set : **A:\PTO.AMC.txt**
Output Set: **N:\CRF4\06272006\I857581C.raw**

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260 ttgttgaggg tgaggtcagc ggggttttcc ttgacacttt gcttgaattc gctgaggatg    900
261 agaccatgga gatcaaaatc accaaggacc acatcgaggg tcttgttgtc gacttttttct    960
262 cggcaggaac agactccaca gcggtggcaa cagagtgggc attggcagaa ctcatcaaca   1020
263 atcctaaggt gttggaaaag gctcgtgagg aggtctacag tgttgtggga aaggacagac   1080
264 ttgtggacga agttgacact caaaaccttc cttacattag agcaatcgtg aaggagacat   1140
265 tccgcatgca cccgccactc ccagtggtca aaagaaagtg cacagaagag tgtgagatta   1200
266 atggatatgt gatcccagag ggagcattga ttctcttcaa tgtatggcaa gtaggaagag   1260
267 accccaaata ctgggacaga ccatcggagt tccgtcctga gaggttccta gagacagggg   1320
268 ctgaagggga agcagggcct cttgatctta ggggacaaca ttttcaactt ctcccatttg   1380
269 ggtctgggag gagaatgtgc cctggagtca atctggctac ttcgggaatg gcaacacttc   1440
270 ttgcatctct tattcagtgc ttcgacttgc aagtgctggg tccacaagga cagatattga   1500
271 agggtggtga cgccaaagtt agcatggaag agagagccgg cctcactgtt ccaagggcac   1560
272 atagtcttgt ctgtgttcca cttgcaagga tcggcgttgc atctaaactc ctttcttaat   1620
273 taagatcatc atcatatata atatttactt tttgtgtgtt gataatcatc atttcaataa   1680
274 ggtctcgttc atctactttt tatgaagtat ataagccctt ccatgcacat tgtatcatct   1740
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280 <212> TYPE: PRT
281 <213> ORGANISM: Glycine max
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288             20                  25                  30
290 Pro Asn Pro Pro Ser Pro Lys Pro Arg Leu Pro Phe Ile Gly His Leu
291         35                  40                  45
293 His Leu Leu Lys Asp Lys Leu Leu His Tyr Ala Leu Ile Asp Leu Ser
294     50                  55                  60
296 Lys Lys His Gly Pro Leu Phe Ser Leu Tyr Phe Gly Ser Met Pro Thr
297 65                  70                  75                  80
299 Val Val Ala Ser Thr Pro Glu Leu Phe Lys Leu Phe Leu Gln Thr His
300                 85                  90                  95
302 Glu Ala Thr Ser Phe Asn Thr Arg Phe Gln Thr Ser Ala Ile Arg Arg
303             100                 105                 110
305 Leu Thr Tyr Asp Ser Ser Val Ala Met Val Pro Phe Gly Pro Tyr Trp
306         115                 120                 125
308 Lys Phe Val Arg Lys Leu Ile Met Asn Asp Leu Pro Asn Ala Thr Thr
309     130                 135                 140
311 Val Asn Lys Leu Arg Pro Leu Arg Thr Gln Gln Thr Arg Lys Phe Leu
312 145                 150                 155                 160
314 Arg Val Met Ala Gln Gly Ala Glu Ala Gln Lys Pro Leu Asp Leu Thr
315                 165                 170                 175
317 Glu Glu Leu Leu Lys Trp Thr Asn Ser Thr Ile Ser Met Met Met Leu
318             180                 185                 190
320 Gly Glu Ala Glu Glu Ile Arg Asp Ile Ala Arg Glu Val Leu Lys Ile
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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 06/27/2006
PATENT APPLICATION: **US/09/857,581C** TIME: 16:52:48

Input Set : **A:\PTO.AMC.txt**
Output Set: **N:\CRF4\06272006\I857581C.raw**

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:66; Xaa Pos. 10,16,23,25,39,48,60,73,74,95,96,102,110,112,117,118,121
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Seq#:66; Xaa Pos. 223,253,259,263,264,268,272,285,292,293,294,301,306,311
Seq#:66; Xaa Pos. 312,325,328,329,334,342,377,381,385,387,393,394,402,404
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VERIFICATION SUMMARY DATE: 06/27/2006
PATENT APPLICATION: **US/09/857,581C** TIME: 16:52:48

Input Set : **A:\PTO.AMC.txt**
Output Set: **N:\CRF4\06272006\I857581C.raw**

L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:3460 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66 after pos.:0
M:341 Repeated in SeqNo=66

Raw Sequence Listing before editing (for reference only)

IFW16

RAW SEQUENCE LISTING DATE: 06/27/2006
PATENT APPLICATION: US/09/857,581C TIME: 16:15:25

Input Set : A:\corrected seq lst 6-19-06.txt
Output Set: N:\CRF4\06272006\I857581C.raw

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  3 <110> APPLICANT: Fader, Gary M.
  4       Jung, Woosuk
  5       Brian, McGonigle
  6       Odell, Joan T.
  7       Yu, Xiaodan
  9 <120> TITLE OF INVENTION: Nucleic Acid Fragments Encoding Isoflavone Synthase
 11 <130> FILE REFERENCE: BB1339RCE
 13 <140> CURRENT APPLICATION NUMBER: 09/857,581C
C--> 14 <141> CURRENT FILING DATE: 2001-06-05
 16 <150> PRIOR APPLICATION NUMBER: PCT/US00/01,772
 17 <151> PRIOR FILING DATE: 2000-01-26
 19 <150> PRIOR APPLICATION NUMBER: 60/117,769
 20 <151> PRIOR FILING DATE: 1999-01-27
 22 <150> PRIOR APPLICATION NUMBER: 60/144,783
 23 <151> PRIOR FILING DATE: 1990-07-20
 25 <150> PRIOR APPLICATION NUMBER: 60/156,094
 26 <151> PRIOR FILING DATE: 1999-09-24
 28 <160> NUMBER OF SEQ ID NOS: 66
 30 <170> SOFTWARE: PatentIn version 3.3
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Does Not Comply
Corrected Diskette Needed

P.8

ERRORED SEQUENCES

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3110 <210> SEQ ID NO: 66
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3113 <213> ORGANISM: Artificial Sequence
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RAW SEQUENCE LISTING DATE: 06/27/2006
PATENT APPLICATION: **US/09/857,581C** TIME: 16:15:26

Input Set : **A:\corrected seq lst 6-19-06.txt**
Output Set: **N:\CRF4\06272006\I857581C.raw**

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RAW SEQUENCE LISTING DATE: 06/27/2006
PATENT APPLICATION: **US/09/857,581C** TIME: 16:15:26

Input Set : **A:\corrected seq lst 6-19-06.txt**
Output Set: **N:\CRF4\06272006\I857581C.raw**

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3241 <223> OTHER INFORMATION: Xaa=Gln or Arg
3243 <220> FEATURE:
3244 <221> NAME/KEY: MISC_FEATURE
3245 <222> LOCATION: (175)..(175)
3246 <223> OTHER INFORMATION: Xaa=Val or Leu
3248 <220> FEATURE:
3249 <221> NAME/KEY: MISC_FEATURE
3250 <222> LOCATION: (183)..(183)
3251 <223> OTHER INFORMATION: Xaa=Ala or Thr
3253 <220> FEATURE:
3254 <221> NAME/KEY: MISC_FEATURE
3255 <222> LOCATION: (187)..(187)
3256 <223> OTHER INFORMATION: Xaa=Thr or Ile

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3258 <220> FEATURE:
3259 <221> NAME/KEY: MISC_FEATURE
3260 <222> LOCATION: (191)..(191)
3261 <223> OTHER INFORMATION: Xaa=Met or Val
3263 <220> FEATURE:
3264 <221> NAME/KEY: MISC_FEATURE
3265 <222> LOCATION: (209)..(209)
3266 <223> OTHER INFORMATION: Xaa=Phe or Tyr
3268 <220> FEATURE:
3269 <221> NAME/KEY: MISC_FEATURE
3270 <222> LOCATION: (219)..(219)
3271 <223> OTHER INFORMATION: Xaa=Arg or Trp
3273 <220> FEATURE:
3274 <221> NAME/KEY: MISC_FEATURE
3275 <222> LOCATION: (223)..(223)
3276 <223> OTHER INFORMATION: Xaa=Tyr or His
3278 <220> FEATURE:
3279 <221> NAME/KEY: MISC_FEATURE
3280 <222> LOCATION: (253)..(253)
3281 <223> OTHER INFORMATION: Xaa=Gly or Glu
3283 <220> FEATURE:
3284 <221> NAME/KEY: MISC_FEATURE
3285 <222> LOCATION: (259)..(259)
3286 <223> OTHER INFORMATION: Xaa=Lys or Glu
3288 <220> FEATURE:
3289 <221> NAME/KEY: MISC_FEATURE
3290 <222> LOCATION: (263)..(263)
3291 <223> OTHER INFORMATION: Xaa=Val or Asp
3293 <220> FEATURE:
3294 <221> NAME/KEY: MISC_FEATURE
3295 <222> LOCATION: (264)..(264)
3296 <223> OTHER INFORMATION: Xaa=Val, Asp, or Ile
3298 <220> FEATURE:
3299 <221> NAME/KEY: MISC_FEATURE
3300 <222> LOCATION: (268)..(268)
3301 <223> OTHER INFORMATION: Xaa=Ala or Val
3303 <220> FEATURE:
3304 <221> NAME/KEY: MISC_FEATURE
3305 <222> LOCATION: (272)..(272)
3306 <223> OTHER INFORMATION: Xaa=Phe or Leu
3308 <220> FEATURE:
3309 <221> NAME/KEY: MISC_FEATURE
3310 <222> LOCATION: (285)..(285)
3311 <223> OTHER INFORMATION: Xaa=Thr or Met
3313 <220> FEATURE:
3314 <221> NAME/KEY: MISC_FEATURE
3315 <222> LOCATION: (292)..(292)
3316 <223> OTHER INFORMATION: Xaa=ANY AMINO ACID
3318 <220> FEATURE:
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3319 <221> NAME/KEY: MISC_FEATURE
3320 <222> LOCATION: (293)..(293)
3321 <223> OTHER INFORMATION: Xaa=ANY AMINO ACID
3323 <220> FEATURE:
3324 <221> NAME/KEY: MISC_FEATURE
3325 <222> LOCATION: (294)..(294)
3326 <223> OTHER INFORMATION: Xaa=Thr or Ile
3328 <220> FEATURE:
3329 <221> NAME/KEY: MISC_FEATURE
3330 <222> LOCATION: (301)..(301)
3331 <223> OTHER INFORMATION: Xaa=Phe or Leu
3333 <220> FEATURE:
3334 <221> NAME/KEY: MISC_FEATURE
3335 <222> LOCATION: (306)..(306)
3336 <223> OTHER INFORMATION: Xaa=Thr or Ile
3338 <220> FEATURE:
3339 <221> NAME/KEY: MISC_FEATURE
3340 <222> LOCATION: (311)..(311)
3341 <223> OTHER INFORMATION: Xaa=Val or Glu
3343 <220> FEATURE:
3344 <221> NAME/KEY: MISC_FEATURE
3345 <222> LOCATION: (312)..(312)
3346 <223> OTHER INFORMATION: Xaa=Val or Ala
3348 <220> FEATURE:
3349 <221> NAME/KEY: MISC_FEATURE
3350 <222> LOCATION: (325)..(325)
3351 <223> OTHER INFORMATION: Xaa=Arg or Lys
3353 <220> FEATURE:
3354 <221> NAME/KEY: MISC_FEATURE
3355 <222> LOCATION: (328)..(328)
3356 <223> OTHER INFORMATION: Xaa=Gln or Glu
3358 <220> FEATURE:
3359 <221> NAME/KEY: MISC_FEATURE
3360 <222> LOCATION: (329)..(329)
3361 <223> OTHER INFORMATION: Xaa=ANY AMINO ACID
3363 <220> FEATURE:
3364 <221> NAME/KEY: MISC_FEATURE
3365 <222> LOCATION: (334)..(334)
3366 <223> OTHER INFORMATION: Xaa=Val or Ala
3368 <220> FEATURE:
3369 <221> NAME/KEY: MISC_FEATURE
3370 <222> LOCATION: (342)..(342)
3371 <223> OTHER INFORMATION: Xaa=Arg or Ile
3373 <220> FEATURE:
3374 <221> NAME/KEY: MISC_FEATURE
3375 <222> LOCATION: (377)..(377)
3376 <223> OTHER INFORMATION: Xaa=Thr or Ile
3378 <220> FEATURE:
3379 <221> NAME/KEY: MISC_FEATURE

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3380 <222> LOCATION: (381)..(381)
3381 <223> OTHER INFORMATION: Xaa=Glu or Gly
3383 <220> FEATURE:
3384 <221> NAME/KEY: MISC_FEATURE
3385 <222> LOCATION: (385)..(385)
3386 <223> OTHER INFORMATION: Xaa=Tyr, His, or Cys
3388 <220> FEATURE:
3389 <221> NAME/KEY: MISC_FEATURE
3390 <222> LOCATION: (387)..(387)
3391 <223> OTHER INFORMATION: Xaa=Ile or Thr
3393 <220> FEATURE:
3394 <221> NAME/KEY: MISC_FEATURE
3395 <222> LOCATION: (393)..(393)
3396 <223> OTHER INFORMATION: Xaa=Val or Ile
3398 <220> FEATURE:
3399 <221> NAME/KEY: MISC_FEATURE
3400 <222> LOCATION: (394)..(394)
3401 <223> OTHER INFORMATION: Xaa=Leu or Pro
3403 <220> FEATURE:
3404 <221> NAME/KEY: MISC_FEATURE
3405 <222> LOCATION: (402)..(402)
3406 <223> OTHER INFORMATION: Xaa=Arg or Lys
3408 <220> FEATURE:
3409 <221> NAME/KEY: MISC_FEATURE
3410 <222> LOCATION: (404)..(404)
3411 <223> OTHER INFORMATION: Xaa=Ser or Pro
3413 <220> FEATURE:
3414 <221> NAME/KEY: MISC_FEATURE
3415 <222> LOCATION: (413)..(413)
3416 <223> OTHER INFORMATION: Xaa=Ser or Phe
3418 <220> FEATURE:
3419 <221> NAME/KEY: MISC_FEATURE
3420 <222> LOCATION: (422)..(422)
3421 <223> OTHER INFORMATION: Xaa=Glu or Gly
3423 <220> FEATURE:
3424 <221> NAME/KEY: MISC_FEATURE
3425 <222> LOCATION: (428)..(428)
3426 <223> OTHER INFORMATION: Xaa=Gly or Arg
3428 <220> FEATURE:
3429 <221> NAME/KEY: MISC_FEATURE
3430 <222> LOCATION: (429)..(429)
3431 <223> OTHER INFORMATION: Xaa=Pro or Leu
3433 <220> FEATURE:
3434 <221> NAME/KEY: MISC_FEATURE
3435 <222> LOCATION: (435)..(435)
3436 <223> OTHER INFORMATION: Xaa=Gln or Arg
3438 <220> FEATURE:
3439 <221> NAME/KEY: MISC_FEATURE
3440 <222> LOCATION: (447)..(447)

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Input Set : **A:\corrected seq lst 6-19-06.txt**
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3441 <223> OTHER INFORMATION: Xaa=Arg or Gly
3443 <220> FEATURE:
3444 <221> NAME/KEY: MISC_FEATURE
3445 <222> LOCATION: (453)..(453)
3446 <223> OTHER INFORMATION: Xaa=Asn, Ser, or Ile
3448 <220> FEATURE:
3449 <221> NAME/KEY: MISC_FEATURE
3450 <222> LOCATION: (459)..(459)
3451 <223> OTHER INFORMATION: Xaa=Met or Thr
3453 <220> FEATURE:
3454 <221> NAME/KEY: MISC_FEATURE
3455 <222> LOCATION: (485)..(485)
3456 <223> OTHER INFORMATION: Xaa=Asp or Gly
3458 <400> SEQUENCE: 66
W--> 3460 Met Leu Leu Glu Leu Ala Leu Gly Leu Xaa Val Leu Ala Leu Phe Xaa
3461 1               5                  10                  15
3463 His Leu Arg Pro Thr Pro Xaa Ala Xaa Ser Lys Ala Leu Arg His Leu
3464             20                  25                  30
3466 Pro Asn Pro Pro Ser Pro Xaa Pro Arg Leu Pro Phe Ile Gly His Xaa
3467         35                  40                  45
3469 His Leu Leu Lys Asp Lys Leu Leu His Tyr Ala Xaa Ile Asp Leu Ser
3470     50                  55                  60
3472 Lys Lys His Gly Pro Leu Phe Ser Xaa Xaa Phe Gly Ser Met Pro Thr
3473 65                  70                  75                  80
3475 Val Val Ala Ser Thr Pro Glu Leu Phe Lys Leu Phe Leu Gln Xaa Xaa
3476                 85                  90                  95
3478 Glu Ala Thr Ser Phe Xaa Thr Arg Phe Gln Thr Ser Ala Xaa Arg Xaa
3479             100                 105                 110
3481 Leu Thr Tyr Asp Xaa Xaa Val Ala Xaa Xaa Pro Xaa Gly Pro Tyr Trp
3482         115                 120                 125
3484 Xaa Phe Val Arg Lys Leu Ile Met Asn Asp Leu Leu Asn Ala Thr Thr
3485     130                 135                 140
3487 Val Asn Xaa Leu Arg Pro Leu Arg Thr Gln Gln Ile Arg Lys Xaa Leu
3488 145                 150                 155                 160
3490 Arg Xaa Met Ala Gln Xaa Ala Glu Ala Xaa Lys Pro Leu Asp Xaa Thr
3491                 165                 170                 175
3493 Glu Glu Leu Leu Lys Trp Xaa Asn Ser Thr Xaa Ser Met Met Xaa Leu
3494             180                 185                 190
3496 Gly Glu Ala Glu Glu Ile Arg Asp Ile Ala Arg Glu Val Leu Lys Ile
3497         195                 200                 205
3499 Xaa Gly Glu Tyr Ser Leu Thr Asp Phe Ile Xaa Pro Leu Lys Xaa Leu
3500     210                 215                 220
3502 Lys Val Gly Lys Tyr Glu Lys Arg Ile Asp Asp Ile Leu Asn Lys Phe
3503 225                 230                 235                 240
3505 Asp Pro Val Val Glu Arg Val Ile Lys Lys Arg Arg Xaa Ile Val Arg
3506                 245                 250                 255
3508 Arg Arg Xaa Asn Gly Glu Xaa Xaa Glu Gly Glu Xaa Ser Gly Val Xaa
3509             260                 265                 270
3511 Leu Asp Thr Leu Leu Glu Phe Ala Glu Asp Glu Thr Xaa Glu Ile Lys
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3512          275                 280                 285
3514 Ile Thr Lys Xaa Xaa Xaa Lys Gly Leu Val Val Asp Xaa Phe Ser Ala
3515     290                 295                 300
3517 Gly Xaa Asp Ser Thr Ala Xaa Xaa Thr Glu Trp Ala Leu Ala Glu Leu
3518 305                 310                 315                 320
3520 Ile Asn Asn Pro Xaa Val Leu Xaa Xaa Ala Arg Glu Glu Xaa Tyr Ser
3521                 325                 330                 335
3523 Val Val Gly Lys Asp Xaa Leu Val Asp Glu Val Asp Thr Gln Asn Leu
3524             340                 345                 350
3526 Pro Tyr Ile Arg Ala Ile Val Lys Glu Thr Phe Arg Met His Pro Pro
3527         355                 360                 365
3529 Leu Pro Val Val Lys Arg Lys Cys Xaa Glu Glu Cys Xaa Ile Asn Gly
3530     370                 375                 380
3532 Xaa Val Xaa Pro Glu Gly Ala Leu Xaa Xaa Phe Asn Val Trp Gln Val
3533 385                 390                 395                 400
3535 Gly Xaa Asp Xaa Lys Tyr Trp Asp Arg Pro Ser Glu Xaa Arg Pro Glu
3536                 405                 410                 415
3538 Arg Phe Leu Glu Thr Xaa Ala Glu Gly Glu Ala Xaa Xaa Leu Asp Leu
3539             420                 425                 430
3541 Arg Gly Xaa His Phe Gln Leu Leu Pro Phe Gly Ser Gly Arg Xaa Met
3542         435                 440                 445
3544 Cys Pro Gly Val Xaa Leu Ala Thr Ser Gly Xaa Ala Thr Leu Leu Ala
3545     450                 455                 460
3547 Ser Leu Ile Gln Cys Phe Asp Leu Gln Val Leu Gly Pro Gln Gly Gln
3548 465                 470                 475                 480
3550 Ile Leu Lys Gly Xaa Asp Ala Lys Val Ser Met Glu Glu Arg Ala Gly
3551                 485                 490                 495
3553 Leu Thr Val Pro Arg Ala His Ser Leu Val Cys Val Pro Leu Ala Arg
3554             500                 505                 510
3556 Ile Gly Val Ala Ser Lys Leu Leu Ser
3557         515                 520
E--> 3560 59
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L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:3460 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66 after pos.:0
M:341 Repeated in SeqNo=66
L:3560 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:66